XP-002090820

Rec'd 22-01-98
Complete = (8)

Hsaj03147.R55u031

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NI   e1246029
DT   22-JAN-1998 (Rel. 54, Created)
DT   03-JUL-1998 (Rel. 56, Last updated, Version 2)
DE   Homo sapiens complete genomic sequence between D16S3070 and
DE   D16S3275, containing Familial Mediterranean Fever gene disease
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KW   HUMNK4 gene; mareno gene; marenostrin; metalloproteinase; mmp20 gene;
KW   olfactory receptor; zinc finger protein; znfmf gene.
OS   Homo sapiens (human)
OC   Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates;
OC   Catarrhini; Hominidae; Homo.
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RA   Bernot A., Heilig R., Clepet C., Smaoui N., da Silva C., Petit J.L.,
RA   Devaud C., Chiannilkulchai N., Fizames C., Samson D., Cruaud C.,
RA   Caloustian C., Gyapay G., Delpech M., Weissenbach J.;
RT   "A transcriptional map of the FMF region";
RL   Genomics 50:147-160(1998).
RN   [2]
RP   1-239566
RA   Bernot A.;
RT   ;
RL   Submitted (07-JAN-1998) to the EMBL/GenBank/DDBJ databases.
RL   GENOSCOPE - Centre National de Sequencage, 2 rue Gaston Cremieux, EVRY
RL   BP191, FRANCE.
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214751 TGTTTTTTCA ATAGCTGAAA ATACAGGTGT GTGCTACCAC GCCTGGCTCA

214801 TTTTTTTGTA TTTTTGGTAG AGACAGGGTT TCACCATGTT GCCCAAGCTG

214851 GTCTCAAACT CCTGAGCTCT AGCAATCCAC CCACCTTAGC CTCCCAAAGT

214901 GCTGAGATTA TTTATAAGTG TGAGCCACCA TGCCTAGCCT ATATTTCTCA

214951 AATAGTACTT TTCAAACGTT TTAAGCAGGA CAACTGTTTC AAATAAGATC

215001 TTACACAGGT CAGCAAAACC CAGATTATAC GCTTTAAAAA TAAAAATAAG

215051 CCGGGCACGG TGGCCTATGT CTGAAATCCC AGCACTTTGG GAGGCTGAGG

215101 CGGGAGGATC ACTTGAGGTC AGGAGTTCGA GACCAGCCTG GCCAACATGG

215151 TGAACAGCAC TGAGAAGATT CTATACCAAC CCCGTTCATA TGATTGCATA

215201 GCAATCCCTT TTTGCTAACC TAGAGATGTT TGTCTGACAT GAGTATTAAT

215251 CATAAATGTA GTGAAGAAGT CTCAAAGAAC AGGTGTTCCA GCTCCTGCCT

215301 TTCGTGACCC TGACCTGCTT CTTAAAAAAC CCATTAGGAG CCTGAAGGAA

215351 GTTTACTAAC CCAGTTCCAA AGGCCAGAGT GAAGAAAGGG ACGTTCCTGA

215401 ACTAAAGTCA TCTGGATTTT GGTAGACCTG AGACTCCCAA TCCCCAGGTC

215451 AGAGTGAGCT GCTCTGAGCT CCTGGTCCCC TTTCCCACAA AGCAGCCAGC

215501 ACTCAGCACT GGATGAGGAG GAGGCCTGGG CCCGCTTACC CTGAATGGCT

215551 GCCCTGTGGA GCTCCTCGGC CAGCAGGCGC TGGTTGATGG CCCGCAGGAC

215601 CTGCAGGGTG AGCTGCACGG CGTACTCTTC CCCATAGTAG GTGACCAGCA

215651 GAGTGGCCAT CTTCACCGGC CTGGCTCTCT GGATCTGGCT CCGGGGGATC

215701 CTGGAGTGCT CCTTCTGCAC ACTGGTGTTC TGCAGCTTGA ACTTGAACTT

215751 CTCGAAGTCA TAGGGCACCA GCTCCTCCAG GGTGGACAGC AGATGGTCAC

215801 TAGGGGTCTT AGCCATGGTG CTGAGCAGGA GAGGCTCGAG CCAGCTGTCT

215851 GGCTTCTGGT AGGAAAAGAA GCCTCTGTCC TTGGTGAGCA AGAAAAGGCA

215901 GGTTGTGAAA TAGCGGAAAA GGCACAGGAA ATGCTCTGTG TCTTGGTGGG